

## Inducible Cassette Exchange: A Rapid and Efficient System Enabling Conditional Gene Expression in Embryonic Stem and Primary Cells

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### ABSTRACT

Genetic modification is critically enabling for studies addressing specification and maintenance of cell fate; however, methods for engineering modifications are inefficient. We demonstrate a rapid and efficient recombination system in which an inducible, floxed *cre* allele replaces itself with an incoming transgene. We target this inducible cassette exchange (ICE) allele to the (*HPRT*) locus and demonstrate recombination in murine embryonic stem cells (ESCs) and primary cells from derivative ICE mice. Using lentivectors, we demonstrate

recombination at a randomly integrated ICE locus in human ESCs. To illustrate the utility of this system, we insert the myogenic regulator, *Myf5*, into the ICE locus in each platform. This enables efficient directed differentiation of mouse and human ESCs into skeletal muscle and conditional myogenic transdetermination of primary cells cultured in vitro. This versatile tool is thus well suited to gain-of-function studies probing gene function in the specification and reprogramming of cell fate. *STEM CELLS* 2011;29:1580–1588

Disclosure of potential conflicts of interest is found at the end of this article.

### INTRODUCTION

The isolation of mouse embryonic stem cells (mESCs) and human ESCs (hESCs) from blastocyst embryos [1–3] has enabled the modeling of early embryonic developmental events and new strategies for tissue engineering. Many studies, especially those probing specification and maintenance of cell fate and mechanisms of differentiation, require genetic modification and would be facilitated by robust methods to modify cells for conditional gene expression. Because every locus differs in its own unique expressability, particularly the differential level of openness or extent to which it is subject to silencing within different lineages, there are obvious advantages to targeting a known and defined locus, especially when comparing the effects of expressing two different genes. Homologous recombination, commonly performed in mESCs [4, 5] is very difficult to implement in other cell types including hESCs [6], and although routine in the murine system, suffers from low frequencies ( $10^{-6}$ ) of recombination and comparatively high frequencies of random integration. Chimeric zinc finger nucleases [7] dramatically improve rates of homologous recombination but the exposure of the genome to such nucleases may result in off-target mutations [8]. Here we describe a system that enables high-efficiency cassette-exchange replacement of a doxycycline-inducible floxed *cre* transgene

with a new transgene of interest. The system, referred to as inducible cassette exchange (ICE), enables high efficiency integration of genes of interest into cells bearing a single-copy ICE locus. We have created platforms for this recombination system in mESCs, primary cells derived from ICE mice, and hESCs and apply this tool to probe the acquisition and stability of cell fate.

### MATERIALS AND METHODS

#### Generation of A2Lox.cre mESCs

mESCs were cultured on irradiated mouse embryonic fibroblasts (MEFs) in Dulbecco's modified Eagle's medium (DMEM)/15% FBS, penicillin/streptomycin (P/S, Gibco), 2 mM glutamax (Invitrogen, Carlsbad, CA, <http://www.invitrogen.com>), nonessential amino acids, 0.1 mM  $\beta$ -mercaptoethanol, and 100 U/ml leukemia inhibitory factor (LIF) (Peprotech, Rocky Hill, NJ, <http://www.peprotech.com>). For embryoid body (EB) differentiation, ESCs were trypsinized and replated in differentiation medium (IMDM/15% FBS, 200  $\mu$ g/mL transferrin (Sigma, St. Louis, MO, <http://www.sigma-aldrich.com>), 4.5 mM monothioglycerol (Sigma), 50  $\mu$ g/mL ascorbic acid (Sigma), and 2 mM glutamax) for 30 minutes to allow MEFs to adhere. Nonadherent cells ( $10^5$ ) were plated as a cell suspension in low adherence dishes on a slowly

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rotating shaker. To generate A2Lox.cre ESCs, the hypoxanthine-guanine phosphoribosyltransferase (HPRT) 5' repair/targeting plasmid [9] carrying the cassette exchange doxycycline (tetracycline)-responsive promoter (TRE)-2loxP-Δneo inducible target locus [10] was digested with XbaI and ligated to an XbaI-Sall fragment bearing the *cre* transgene from pSalk-*cre* [11]. Sall-linearized DNA (20 μg) was electroporated into 6 × 10<sup>6</sup> A17 mESCs, and selection in ES medium with HAT supplement (Invitrogen) was initiated 24 hours later.

### Generation of inducible GFP and iMyf5 mESC Lines

To generate derivative inducible mESC lines, A2Lox.cre mESC lines were exposed to 500 ng/mL doxycycline for 24 hours, trypsinized, counted, and 2 × 10<sup>5</sup> cells were electroporated (Amaxa, Köln, Germany, <http://www.lonzabio.com> nucleofector 96-well shuttle, solution VPHF-1001, waveform program 96-CG-104) with 4 μg of p2Lox-mMyf5 or p2Lox-EGFP plasmid and plated on neo-MEFs (Specialty Media, St. Louis, MO, <http://www.sigma-aldrich.com>). Selection was initiated in 300 μg/mL G418 (Gibco, Carlsbad, CA, <http://www.invitrogen.com>) 24 hours after plating and maintained for 10 days. Colonies were picked at day 8 and replated on neo-MEFs for expansion.

For myogenic differentiation, iMyf5 mESCs were differentiated as EBs for 2 days, then attached to gelatin-coated plates in EB differentiation medium. In the initial 4 days, cells were cultured in DMEM/10% FBS with 500 ng/mL doxycycline. They were then switched to DMEM/2% horse serum (HS) with 500 ng/mL doxycycline and cultured for an additional 4 days.

### Generation of iDsRed2 and iMyf5 Kidney Cell Lines

ICE mice were derived at the University of Texas UT Southwestern Transgenic Core Facility by blastocyst injection of ZX1 mESCs, an ICE mESC line related to A2Lox.cre but with an improved TRE promoter [12]. Mice were housed in a pathogen-free environment and cared for under the guidance of the University of Texas UT Southwestern and University of Minnesota Institutional Animal Care and Use Committees. Primary kidney cells were obtained from collagenase I-treated, minced whole kidney pieces from male ICE mice. These were allowed to attach in DMEM/10% FBS and P/S at 37°C, enabling constituent cells to spread over the surface of a T75 flask. Cells were then passaged by trypsinization at 80% confluence. p2Lox-DsRed2 and -Myf5 were introduced by electroporation with an Amaxa nucleofector 96-well shuttle (Nucleofector II: VPI-1002 waveform program U-012). For the quantification of recombination, cells were passaged twice and induced overnight with 500 ng/mL doxycycline. For the derivation of iMyf5 kidney cells, 2 days post-nucleofection, selection was initiated in 75 μg/mL G418 for 4 days, then brought to 100 μg/mL, and maintained over a period of 20 days. During selection, cells were passaged at 80% confluence. For myogenic differentiation, iMyf5 primary kidney cells were induced with 500 ng/mL doxycycline and cultured on gelatin-coated plates in myogenic medium (DMEM/20% FBS, 10 ng/mL basic fibroblast growth factor (bFGF), and 10<sup>-7</sup> M dexamethasone) for 8 days. When the culture reached 100% confluency, medium was replaced with DMEM/2% HS with 500 ng/mL doxycycline for an additional 4 days.

### Lentivirus Production

Lentiviral supernatant was produced in 293T cells cultured in DMEM/10% FBS. Lentiviral plasmid constructs, packaging construct pHRS.2deltaR, and envelop plasmid pCMV-VSV-G were cotransfected using FUGENE HD (Roche, Basel, Switzerland, [www.roche-applied-science.com](http://www.roche-applied-science.com)). Medium was changed after 24 hours and the viral supernatant was collected at 48 hours post-transfection. Filtered supernatant (0.45 μm) was spiked with polybrene (4 μg/mL) and applied to cells plated at low density 1 day before infection. Spin infection was performed at 2,000g at 33°C for 90 minutes in a Sorval Legend RT centrifuge. Cells were

incubated for an additional 3 hours at 37°C/5% CO<sub>2</sub> at which time the supernatant was replaced with fresh medium.

### Generation of A2Lox.cre.I hESCs

H9 hESCs were cultured in mTESR (Stemcell Technologies) supplemented with P/S (Gibco) on growth factor-free matrigel (BD Biosciences)-treated dishes and passaged using Accumax (Millipore, Billerica, MA, [www.millipore.com](http://www.millipore.com)) with Rock inhibitor (Enzo Life Sciences, Farmingdale, NY, <http://www.enzolifesciences.com>). Cells were transduced twice with high titer supernatants of Lenti-rtTA [13], a lentivector bearing rtTA under the ubiquitin C promoter (a modified version of FUGW [14] in which rtTA2SM2 from pUHDrtTA2SM2 [15] was introduced downstream of the Ubiquitin promoter) to generate derivative H9:rtTA hESCs. These cells were then transduced with 2Lox.cre-ires-GFP [13] lentivector over a dilution series of supernatant. Cells were pulsed with 500 ng/mL doxycycline, and bulk sorted from a dilution with <5% transduction, to bias for single-copy integration events. Sorted cells were grown to 50% confluence, pulsed with doxycycline, and sorted a second time to remove clones in which the integration was silenced. When these cells reached 50% confluence, they were single cell sorted into 96-well dishes and clonal cell lines were expanded. A split from each clonal cell line was tested for levels of GFP inducibility, and the best clones were tested for recombination. Clone H9-2Lox.cre.I was selected for all following experiments.

### Generation of H9imCherry and H9iMyf5 hESCs

H9-2Lox.cre.I cells were treated with 500 ng/mL doxycycline for 24 hours and harvested, and five batches of 2 × 10<sup>5</sup> cells were nucleofected with 4 μg p2Lox-mCherry, 4 μg p2Lox-DS-RED2, or p2Lox-hMyf5-ires-mCherry with an Amaxa 96-well nucleofector shuttle using VPHF-5003 solution for H9 cells and the hESC waveform protocol, 96-CB-150. For quantification of recombination, cells were passaged twice and then induced overnight with 500 ng/mL doxycycline. To generate imCherry or iMyf5 derivatives, cells were expanded to 50% confluence on a T75 flask and induced with 200 ng/mL doxycycline overnight. The following day, red fluorescent cells were sorted and plated on a six-well dish. Secondary and tertiary enrichment sorts were performed in the same way.

Cells were differentiated by plating 4 × 10<sup>5</sup> cells per milliliter in low-adherence dishes in mTESR medium for 2 days to allow aggregates to form. Medium was then changed to growth factor-free mTESR (Stemcell Technologies, Vancouver, Canada, [www.stemcell.com](http://www.stemcell.com)) supplemented with 10 ng/mL human bone morphogenetic protein 4 (hBMP4), 3 ng/mL hActivinA, and 10 ng/mL human basic fibroblast growth factor (hbFGF) (all from Peprotech) for 4 days. BMP4 and ActivinA were then withdrawn and bFGF was reduced to 5 ng/mL hbFGF for the next 4 days. At this stage, EBs were plated in adherent dishes to allow attachment in myoblast medium: DMEM/F12 supplemented with 20% FBS, P/S, 10 ng/mL hbFGF, 10<sup>-7</sup> M dexamethasone, and 0.1 mM β-mercaptoethanol. After 2 days of attachment, adherent cells, referred to hereafter as mesenchymal cells, were collected by trypsinization and expanded for up to six passages. For myogenic differentiation, mesenchymal cells were maintained in doxycycline from the point of attachment. Terminal differentiation was induced by switching myoblast medium for DMEM/2% HS for 6 days.

### Immunofluorescence and Western Blotting

Cells were cultured on gelatin-coated coverslips, fixed with 4% paraformaldehyde for 20 minutes, permeabilized by 0.3% Triton X-100 for 30 minutes, and blocked by 10% goat serum for 1 hour at room temperature. Primary mouse anti-MyoD (BD Biosciences cat. no. 554130) or mouse anti-sarcomeric myosin (MF20, Developmental Studies Hybridoma Bank, University of Iowa, IA, [dshb.biology.iowa.edu](http://dshb.biology.iowa.edu)) was incubated in phosphate buffer saline (PBS)/2% goat serum at 4°C overnight, followed by secondary

goat anti-mouse Cy3- or Alexa-555-conjugated antibodies (Sigma, and Invitrogen, respectively) at room temperature for 45 minutes. Cells were counterstained using 4',6-diamidino-2-phenylindole (Invitrogen) to visualize nuclei and mounted in "Immu-Mount" (Thermo Scientific, Pittsburgh, PA, [www.thermoscientific.com](http://www.thermoscientific.com)). Images were acquired on an Axio Observer.Z1 microscope (Zeiss, Oberkochen, Germany, [www.zeiss.de](http://www.zeiss.de)). For Western Blotting, cells were cultured on gelatin-coated plates, and protein was extracted by using RIPA buffer. Primary mouse anti-GAPDH (HRP conjugated) was used to detect the specific band of 36 kD. Primary mouse anti-Myf5 (SC-302 clone C-20) and secondary goat anti-rabbit-HRP conjugated were used to detect the Myf5 specific band at 35 kD. Primary mouse anti-MyoD (BD Biosciences c# 554130), mouse anti-myogenin (BD Bioscience, East Rutherford, NJ, <http://www.bd.com> c# 556358), mouse anti-M-cadherin (BD Bioscience c# 611100), mouse anti-myosin heavy chain (MyHC [DSHB F1.652S]), and secondary anti-mouse HRP conjugated were used to detect specific bands of 45, 34, 130, 180 kD, respectively.

### Quantitative Real Time RT-PCR

Total RNA was extracted with Trizol (Invitrogen) and cDNA was generated using 1  $\mu$ g DNase-treated RNA with oligo-dT primers and TermoScript (Invitrogen). PCR was performed using TaqMan Real Time PCR premixtures on a 7500 RT-PCR System from Applied Biosystems. For muscle-related genes, pre-made probes were purchased from Applied Biosystems: Myod1 Mm00440387\_m1, Myf5 Mm00435125\_m1, Myog Mm00446194\_m1, Mck Mm0432556\_m1, Desmin Mm00802455\_m1, m-Cad Mm00483183\_m1, and Gapdh Mm99999915\_g1. For human probes: MYOD1 Hs00159528\_m1, MYF5 Hs00271574\_m1, MYOG Hs01072232\_m1, DESMIN Hs00157258\_m1, and GAPDH Hs99999905\_m1. All reactions were performed at least in triplicate and the data were analyzed by 7500 System Software (Applied Biosystems, Foster City, CA, [www.appliedbiosystems.com](http://www.appliedbiosystems.com)). The difference in threshold cycle number between transcripts of interest and GAPDH was used to determine the abundance of transcript relative to GAPDH ( $\Delta$ -CT method).

### Statistical Analyses

All experiments were done at least three times. Data shown for real time PCR are the mean  $\pm$  SD. Difference between means was compared by the two-tailed Student's *t* test (GraphPad Prism five) and was considered significantly different at  $p < .05$ .

## RESULTS

### Generation of ICE mESCs

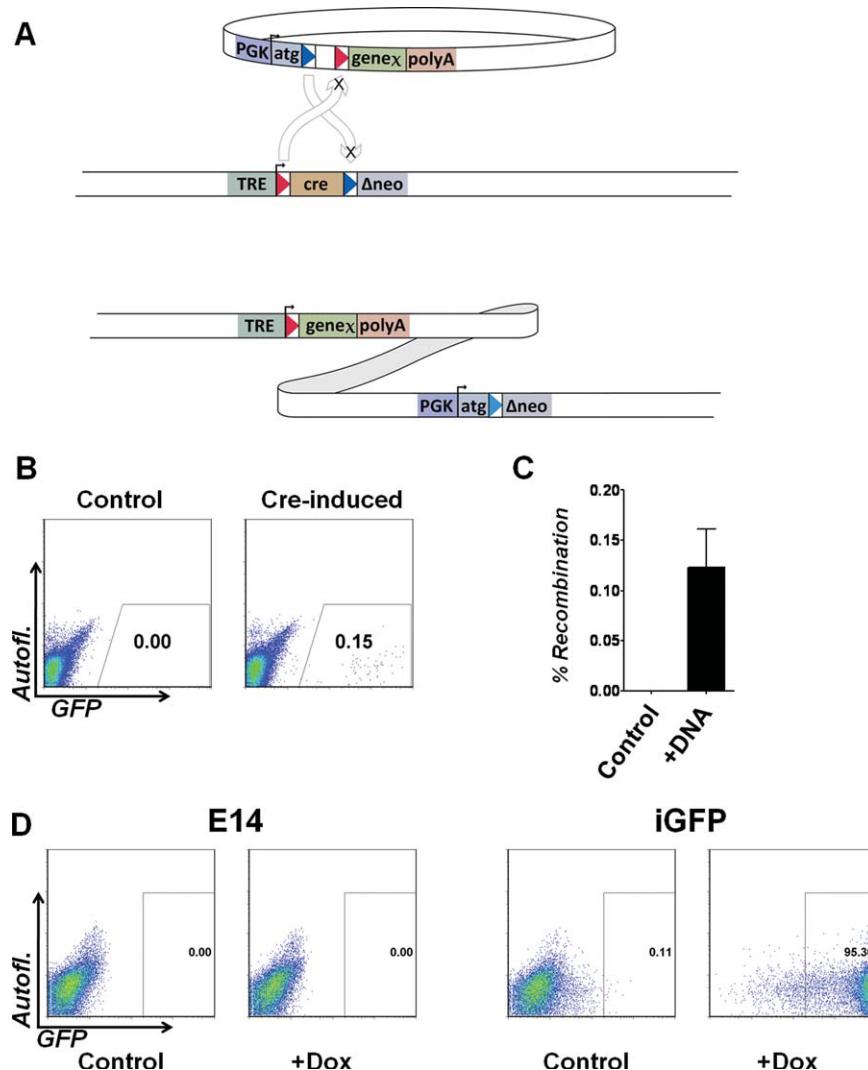
The genomic region upstream of *HPRT* has been demonstrated to be a site at which transgenes express reliably [9, 16-18], presumably due to *HPRT* being a housekeeping gene, active in all cell types and therefore embedded in constitutively open chromatin. We have targeted this site with a doxycycline-inducible *cre* transgene flanked by heterologous, self-incompatible *loxP* sites (Fig. 1A) in A17 mESCs, a derivative of E14Tg2a [19] in which rtTA has been inserted into the *Rosa26* locus [20]. Downstream of the floxed *cre* is *Δneo*, a G418 resistance gene lacking a start codon and promoter [21]. These cells are referred to as A2Lox.cre. Treating cells with doxycycline causes *cre* to be expressed, rendering the cells competent for recombination. When transfected with p2Lox, a plasmid bearing the same heterologous *loxP* sites, a cassette exchange recombination replaces *cre* with a gene of interest from the incoming plasmid. Because the orientation of the *loxP* sites is inverted on the plasmid relative to the chromosome, the plasmid in its entirety (except for the short sequence between the *loxP* sites) is inserted. At the upstream *loxP* site, a gene of interest is brought under the regulation of the doxycycline-inducible promoter, while at the downstream

*loxP* site, a PGK promoter and a start codon are spliced in-frame with the *Δneo* gene. This results in reversion to G418 resistance and selection for precise integration [21].

To measure the efficiency of ICE recombination in mESCs, we treated A2Lox.cre cells with doxycycline for 24 hours to induce *cre* expression or kept untreated cells as controls. We then nucleofected both with p2Lox bearing GFP. After several days of growth in the absence of doxycycline, cells from both arms were treated with doxycycline for 48 hours. In cells nucleofected with p2Lox-GFP but not previously exposed to doxycycline, no integrants were observed; however, in cells previously pulsed with doxycycline to induce *cre*, a recombinant population was observed and the frequency of recombination was determined to be  $>1/1,000$  (the frequency of GFP+ cells, Fig. 1B, 1C). This is approximately three orders of magnitude more efficient than homologous recombination. By subjecting the treated population to G418 selection, individual colonies can be isolated, and these now show inducible expression, both in the ESC stage, as well as when differentiated for several days as EBs (Fig. 1D).

### Directing Mesoderm Differentiation with Myogenic Regulatory Factors

To evaluate the usefulness of this system in probing mechanisms of cell fate determination, we sought to derive a developmental lineage that is poorly represented in conventional mES in vitro differentiation. The skeletal myogenic lineage is very difficult to derive from mESCs, most likely because of a deficiency in the inductive signals that initiate somitogenesis in paraxial mesoderm [22]. To test the ability of this inducible system to direct the lineage choice of differentiating cells, specifically to direct cells into the myogenic lineage, we generated a new inducible mESC line using this system: iMyf5 (for inducible *Myf5*). *Myf5* and its related paralog *MyoD* display significant genetic redundancy in early myogenic development [23]; however, *Myf5* is temporally and genetically upstream of *MyoD* [24, 25], thought to be directly induced by Pax3 in cells of the hypaxial somite [26], and is downregulated following myogenic specification. We differentiated iMyf5 cells in serum suspension (EB) culture for 2 days, conditions under which mesoderm is potently induced. EBs were then allowed to attach, and doxycycline was applied from this time point onward, to induce *Myf5* expression in early mesodermal progenitors as soon as they emerged (Fig. 2A). In the presence of *Myf5* induction, cells with *MyoD*+ nuclei were observed, indicating myogenic commitment, whereas in control cells lacking *Myf5* induction, cells were uniformly *MyoD*-negative (Fig. 2B). After 4 days of monolayer culture, cells were exposed to myogenic differentiation medium (2% HS, conditions that promote myotube formation from myoblasts) for an additional 4 days. In the absence of *Myf5* induction, no cell fusion was observed and MyHC expression was absent; however, in cells exposed to *Myf5*, large numbers of multinucleated MyHC+ myotubes were formed (Fig. 2B). Differentiated cells were evaluated for myogenic gene expression by quantitative RT-PCR (qRT-PCR). As expected, *Myf5* was robustly expressed in doxycycline-treated cells, and cells exposed to *Myf5* for 8 days expressed the downstream myogenic regulatory factors, *MyoD* and *Myogenin*. In addition, various structural gene markers of terminal myogenic development were observed (Fig. 2C, 2D). These data demonstrate the utility of the system for probing the activity of genes that regulate early events in murine embryonic development, in this case showing that *Myf5* expression is sufficient to promote the myogenic commitment of nascent mesoderm and establishing a novel and rapid method of differentiating mESCs into skeletal muscle tissue.

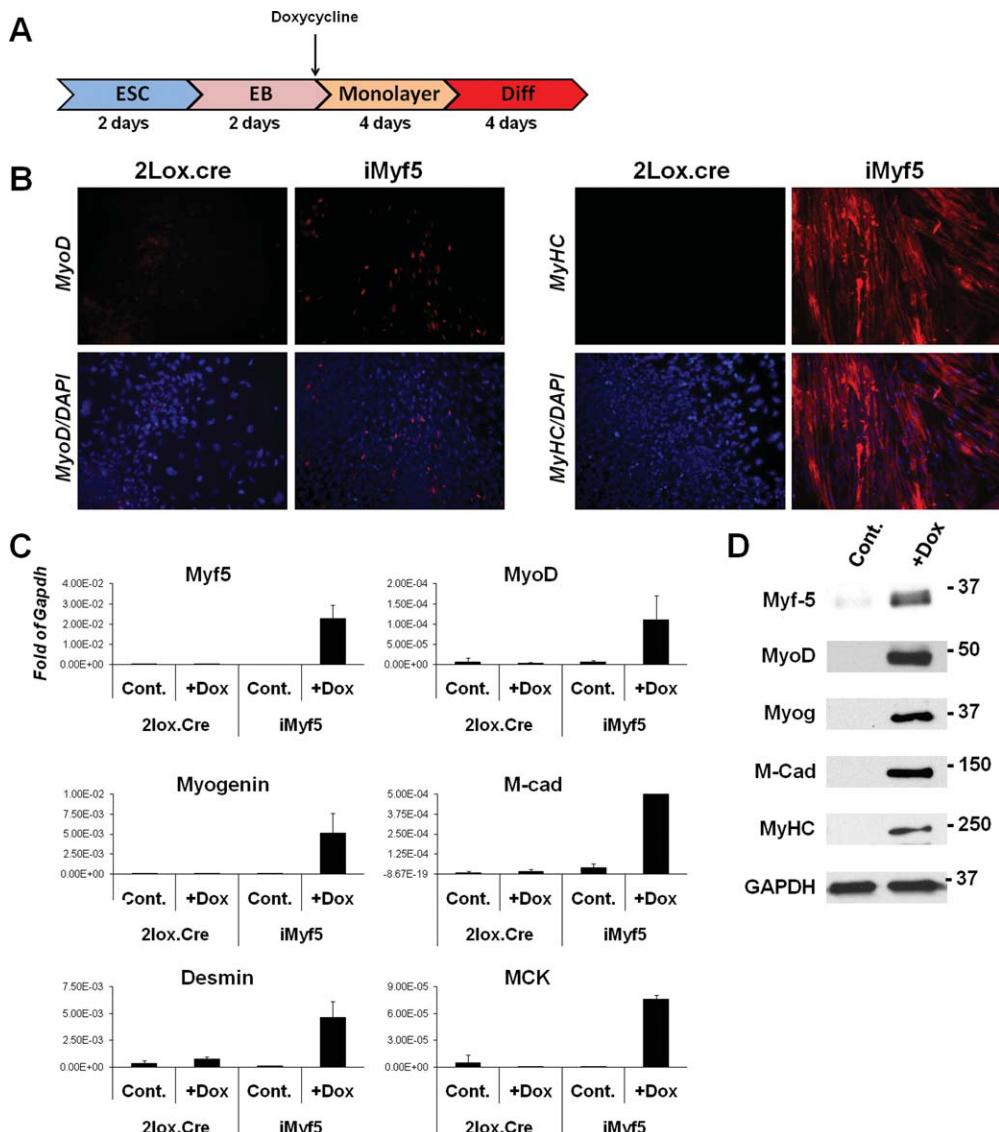


**Figure 1.** Inducible cassette exchange recombination in A2Lox.cre mouse embryonic stem cells (mESCs). (A): The incoming plasmid is shown above the inducible cassette exchange (ICE) target locus, which is integrated at a unique site in the genome (5' of HPRT in the mESCs). At the ICE target locus, *cre* is flanked by heterologous loxP sites and is downstream of a TRE. Arrows indicate recombination between homologous loxP sites. The heterologous loxP sites on the incoming plasmid are in the opposite orientation relative to their cognates on the chromosome, therefore recombination catalyzes the integration of the entire plasmid. *Cre* is exchanged for the incoming gene of interest, rendering that gene doxycycline-inducible in the derivative recombinants. Selection is enabled by correction of the  $\Delta$ neo gene which acquires a PGK promoter, Kozak translational consensus and ATG. (B): FACS Fluorescence activated cell sorting analysis of A2Lox.cre ESCs following recombination in the absence of selection. Cells were either not treated with doxycycline (left) or given a 24-hour pulse of doxycycline (right) prior to nucleofection. After recovery, cells were reinduced with doxycycline to visualize recombinants. (C): Measured frequency of recombination.  $n = 3$ ,  $p < .0049$ . (D): FACS analysis of a derivative inducible GFP cell line and control (E14) exposed to doxycycline. Abbreviations: GFP, green fluorescence protein; PGK, phosphoglycerate kinase; TRE, doxycycline (tetracycline)-responsive promoter.

### ICE Recombination in Human ESCs

To facilitate similar studies in a human system, we sought to derive a hESC with an ICE target locus. We elected to randomly integrate ICE loci and select clones with favorable expression characteristics. We therefore used a modified version of the ICE locus in which an *ires-GFP* reporter follows the *Cre* gene within the domain flanked by loxP sites (Fig. 3A). The entire locus is carried by a lentiviral vector, which allows for efficient introduction of the ICE locus. The presence of the GFP reporter allows the isolation of clonal cell lines that have picked up a genomic integration of the cassette, and their evaluation to determine which have favorable doxycycline responsiveness. In addition, following cassette exchange, recombinants can be distinguished from parental cells

by the absence of inducible GFP expression. We transduced H9 hESCs [1] with this construct at low titer to favor single-copy integration, used a transient pulse of doxycycline to identify integrants, which we sorted and replated. Following expansion of this founder population, we pulsed with doxycycline a second time and single cell sorted to isolate clonal cell lines. We evaluated clones for ICE recombination by nucleofecting each with a plasmid construct bearing mCherry and measuring the frequency of green to red conversions. The best clone, H9.2Lox.cre.I, was karyotyped and selected for further studies. When mCherry was introduced into this clone, a small population of doxycycline-inducible red fluorescent cells among a large background of green fluorescent cells was observed, provided cells had been treated with doxycycline to induce Cre expression just prior to nucleofection. The measured recombination rate was



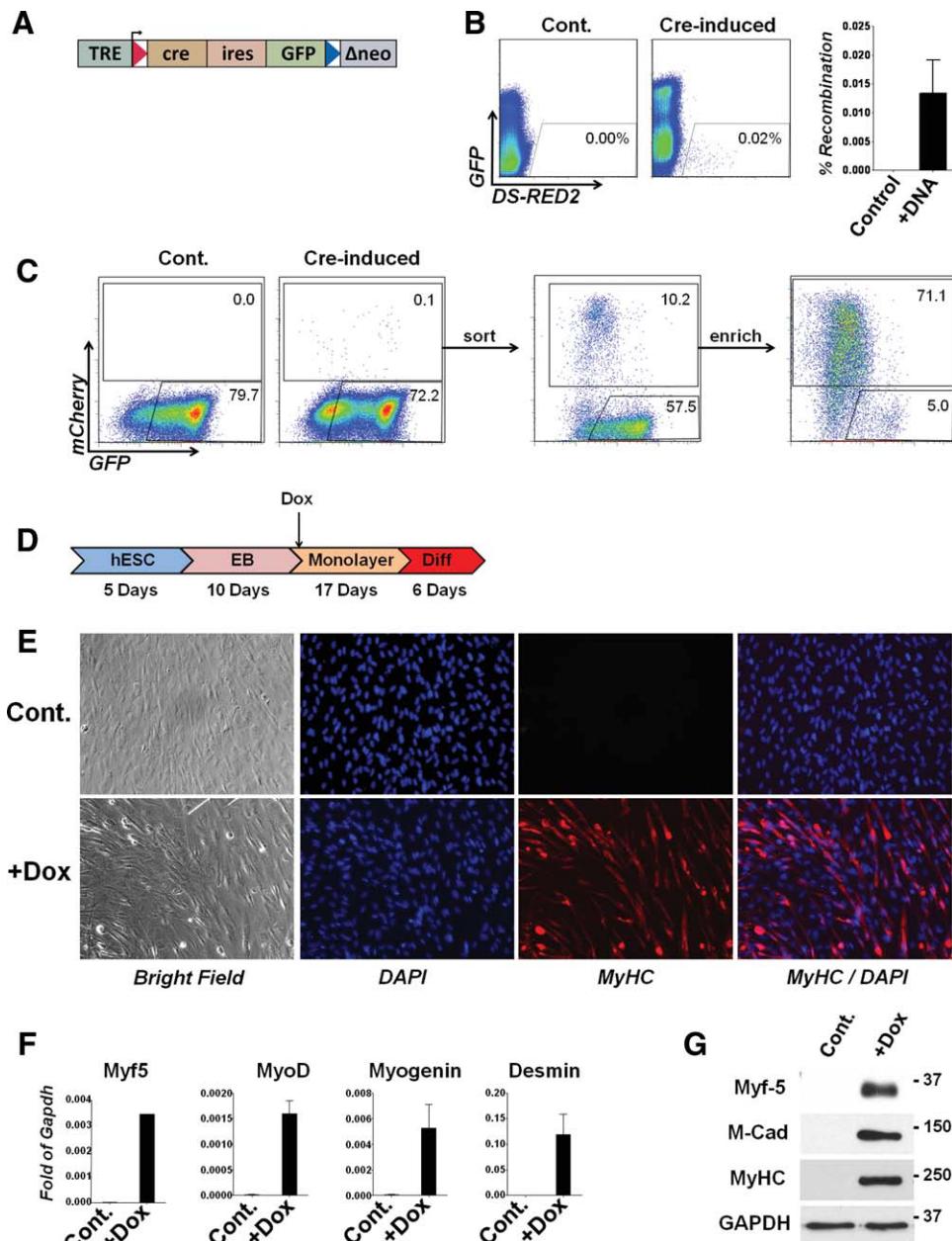
**Figure 2.** Modification of ICE mouse ESC (mESC) with Myf5, and efficient myogenesis in vitro. **(A):** Schematic overview of mouse myogenic differentiation protocol. **(B):** Immunostaining for MyoD or sarcomeric myosin heavy chain (MyHC) in progeny of iMyf5 mESCs or control parental A2Lox.cre mESCs at the final stage of differentiation (in 2% horse serum) in the presence or absence of doxycycline. **(C):** Myogenic gene expression measured by quantitative reverse transcriptase PCR (RT-PCR) in the same cells. **(D):** Myogenic protein expression measured by Western blotting in the same cells. Abbreviations: ESC, embryonic stem cell; EB, embryonic body; DAPI, 4',6-diamidino-2-phenylindole; MCK, muscle creatine kinase; MyHC, myosin heavy chain.

somewhat lower in hESCs than in mouse cells (Fig. 3B). These could be sorted and enriched (Fig. 3C). In cells that were not pretreated with doxycycline to induce Cre expression, nucleofection with p2Lox-mCherry did not promote the emergence of a red fluorescent population. We then introduced *Myf5* into this clone on a plasmid bearing an ires-mCherry reporter and isolated recombinants by flow cytometry. The derivative recombinant cell line was then differentiated in suspension EB culture for 10 days followed by attachment to gelatin-coated dishes in the presence of doxycycline. In both *Myf5*-induced and *Myf5*-uninduced cultures, a mesenchymal cell population emerged from the attached EBs and expanded over the course of 17 days. Medium was then changed to 2% HS to promote differentiation, and at this point a clear difference was observed between the control and induced cultures, with the latter having an abundance of long spindle-shaped cells. After 1 week in differentiation medium, the *Myf5*-induced cultures displayed a substantial amount

of cell fusion, and multinucleated cells were positive for sarcomeric myosin (Fig. 3E). Analysis of gene expression further indicated myogenic differentiation: as seen with the mESC cultures exposed to *Myf5*, the downstream myogenic regulatory factors (MRFs), *MyoD* and *Myogenin*, as well as the differentiation marker, *Desmin* were now coexpressed (Fig. 3F). Markers of myogenic differentiation were also shown by Western blotting (Fig. 3G). These genes were not induced in the control cultures. This efficient and conditional skeletal myogenic differentiation highlights the utility of ICE in hESCs to manipulate genetic pathways controlling lineage specification during differentiation in vitro.

#### ICE Recombination in Primary Cells *In Vitro*

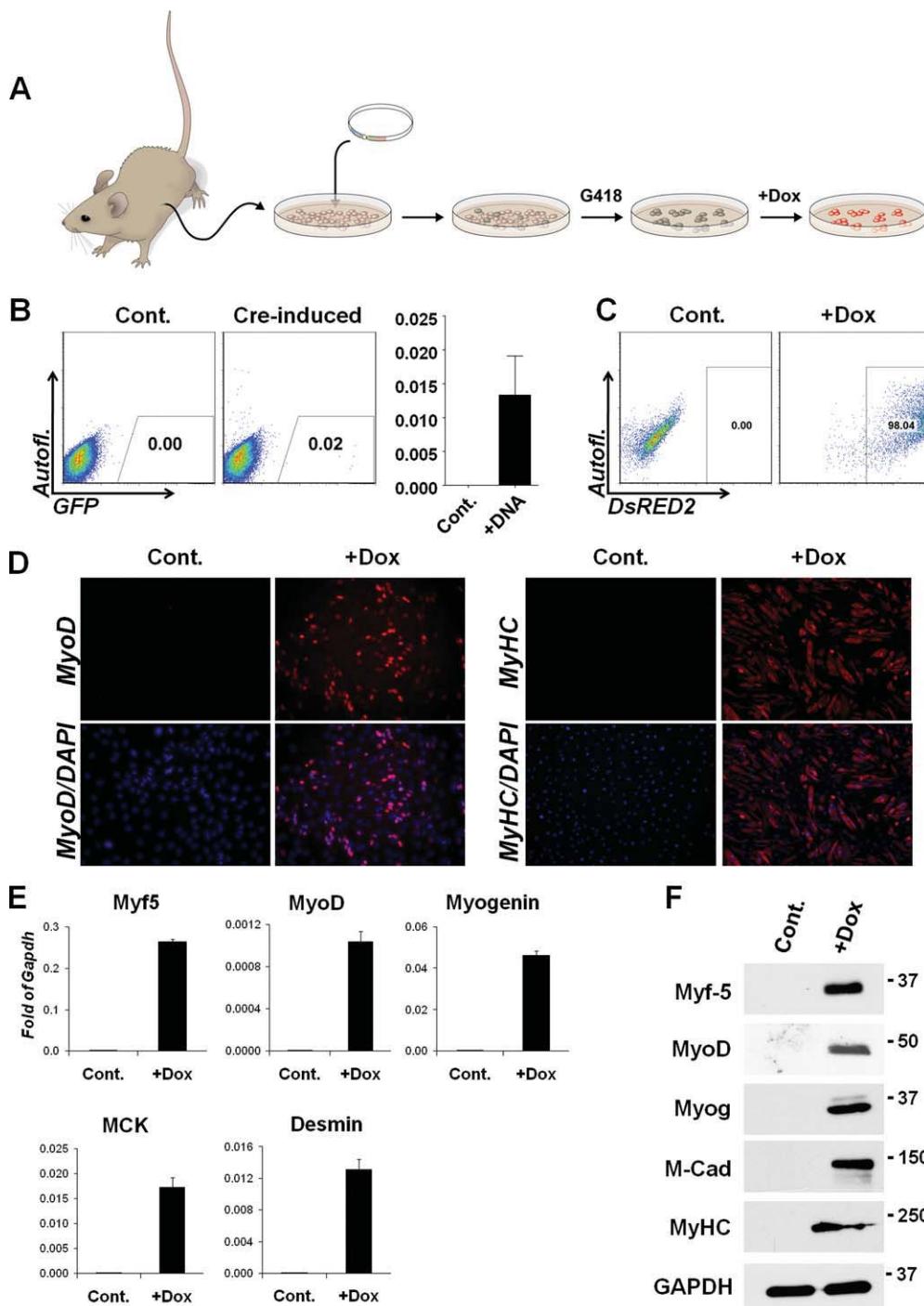
Many applications directed toward probing the stability of cell fate and the feasibility of reprogramming would be facilitated by isogenic methods to compare conditional gene



**Figure 3.** Inducible cassette exchange recombination in H9 hESCs and efficient myogenesis *in vitro*. (A): Schematic of the ICE locus, modified to include an *ires-GFP* reporter gene. (B): Fluorescence activated cell sorting (FACS) analysis of doxycycline-induced H9.2Lox.cre.I ESCs (left panel) and control uninduced H9.2Lox.cre.I cells (right panel) after nucleofection with p2Lox-DsRed2. Inducible red fluorescence replaces inducible green fluorescence in the recombinants. Quantification over several replicate experiments is shown at right. (C): Fluorescent red cells were sorted and enriched. (D): Scheme of human myogenic differentiation. (E): Immunofluorescence for sarcomeric MyHC in H9.iMyf5 cells differentiated in the absence (above) or presence (below) of doxycycline. (F): Myogenic gene expression measured by quantitative RT-PCR in the same cells. (G): Myogenic protein expression measured by Western blotting in the same cells. BF, bright field; DAPI, 4',6-diamidino-2-phenylindole; EB, embryonic body; GAPDH, glyceraldehyde 3-phosphate dehydrogenase; GFP, green fluorescent protein; hECS, human embryonic stem cells; MyHC, myosin heavy chain; TRE, doxycycline (tetracycline)-responsive promoter.

expression in primary cells. However, making multiple modifications of ESCs and taking each cell line through the blastocyst into the germ line to derive new strains would be practical only for a very limited number of constructs. On the other hand, a large number of isogenic modifications could be generated in primary cells from an animal bearing an ICE locus. We therefore derived an ICE mouse strain through blastocyst injection of A2Lox.cre ESCs and tested the efficiency of recombination in primary cells of these mice. In primary cells derived from the kidneys of ICE mice, the recombination

rate, measured by acquisition of doxycycline-inducible DsRED2 protein, was observed to be slightly over 1 in  $10^4$  cells (Fig. 4B). We then integrated *Myf5* and selected a population of G418-resistant cells. Primary cells from the kidney are not myogenic, do not express MyoD, and will not fuse under conditions of growth factor withdrawal (Fig. 4D, control). However, after 1 week of exposure to *Myf5*, myogenic potential was acquired by these cells, indicated by nuclear staining for MyoD, and the ability to fuse to form MyHC-positive myotubes in conventional myoblast differentiation



**Figure 4.** Inducible cassette exchange (ICE) mice and respecification of cell fate in primary cells. **(A):** Schematic overview of recombination in ICE mice: primary cells are expanded and transfected with p2Lox. Recombinants (indicated in dark beige) are selected with G418 and expanded. When doxycycline is applied, the gene brought in on p2Lox (*DsRed2* in the examples shown in **(B)** and **(C)**, below) is induced. **(B):** Recombination in primary kidney cells nucleofected with p2Lox-DsRed2. Controls were not pretreated with doxycycline to induce *cre* expression prior to nucleofection. After nucleofection, cells were allowed to recover and expand in the absence of doxycycline and were later treated with 500 ng/mL doxycycline to visualize recombinants. The histogram at right indicates the average recombination rate over  $n = 3$  independent experiments.  $p = .0161$ . **(C):** Fluorescence activated cell sorting (FACS) for *DsRed2* in the cells shown in **(B)** after growing out in 150  $\mu$ g/mL G418. **(D):** Immunohistochemistry for *MyoD* and *MyHC* in *Myf5*-recombinant kidney cells cultured in 2% horse serum. **(E):** Gene expression analysis by quantitative reverse transcriptase (RT)-PCR for skeletal myogenic markers. **(F):** Myogenic protein expression measured by Western Blotting in the same cells. Abbreviations: DAPI, 4',6-diamidino-2-phenylindole; GADPH, glyceraldehyde 3-phosphate dehydrogenase; GFP, green fluorescent protein; MCK, muscle creatine kinase; MyHC, myosin heavy chain.

medium (Fig. 4D, doxycycline). Transcriptional and protein analysis of skeletal muscle-specific markers including *MyoD*, *Myogenin*, *MCK* (muscle creatine kinase), *Desmin*, and

*M-cadherin*, (Fig. 4E, 4F) further validate the transdetermination of these kidney-derived cells into the skeletal myogenic lineage on induction of *Myf-5*.

## DISCUSSION

The ICE recombination system is versatile—easily set up in any cellular model system, robust—a *cre* expression plasmid is not required, *cre* is expressed at the same level by every cell prior to recombination, and the reaction is self-limiting because after recombination *cre* is gone (indeed, recombination occurs at levels two or three orders of magnitude above conventional homologous recombination), and rapid—a pure population of derivative inducible gene-expressing recombinants can be obtained within a time frame measured in days. This latter feature is particularly valuable in cells of limited lifespan, such as primary cells, or cells in which high passage numbers are undesirable, such as ESCs.

Whether at a known location, such as upstream of *HPRT*, or an unknown locus, such as a proviral integration site, all *cre*-mediated integrations into a given cell type are targeted to the same locus. This reduces variation in expression associated with integration site and copy number differences and eliminates the likelihood that unwanted genetic changes, such as activation of oncogenes, will occur. These features allow elements of an allelic series, for example, a set of different point mutations or deletions within a protein, to be compared to one another directly and phenotypes to be interpreted with confidence.

The conditional doxycycline-inducible nature of the target locus is well suited to gain-of-function experiments. Here, we have focused on determining the effect of a particular transcription factor on the selection of differentiation pathways, or on the acquisition of a novel differentiation pathway. In both the murine and the human system, expression of *Myf5* in early mesodermal progenitor cells is an efficient way of promoting the skeletal myogenic lineage. This lineage is particularly difficult to derive from both mESCs and hESCs—although a few examples of skeletal myogenic differentiation of wild-type ESCs or iPSCs have been reported, these involve

complex protocols based on sorting of rare cells [27, 28]. For studies requiring large quantities of early embryonic myogenic progenitors, genetic modification with inducible expression of genes such as *Myf5*, or its upstream regulators *Pax3* and *Pax7* [22, 29, 30], are ideal.

Additional applications, for example, the modification of iPSC lines or the integration of reporter genes at this locus with minimal position effect are also possible.

## CONCLUSION

The ability of ESCs to differentiate into all somatic cell types and the inherent scalability of cultures initiated with immortal cells makes this a powerful system in which to combine discovery of cellular or developmental phenotypes with mechanistic investigations of the role of a developmental regulatory factors.

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## DISCLOSURE OF POTENTIAL CONFLICTS OF INTEREST

The authors indicate no potential conflicts of interest.

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